

# Discussion of Likelihood Inference for Models with Unobservables: Another View

Thomas A. Louis

## 1. INTRODUCTION

Lee and Nelder identify important issues and provide excellent advice and warnings associated with inferences and interpretations for models with unobserved, latent variables (random effects). Their discussion of prediction versus estimation goals is insightful and I have some sympathy with their call for use of comprehensive probability models. They provide a clear explanation of their h-likelihood approach and a spirited promotion of it. Unfortunately, the value and impact of their advice are compromised by their singular focus on promoting h-likelihood. Their claim that it is an almost universally preferred approach is, to put it mildly, a stretch. The h-likelihood approach by no means “trumps” all competitors and has its own deficits. Over promotion makes the article more of an opinion-piece than a scientific comparison of approaches.

## 2. POINT/COUNTERPOINT

I identify and discuss principal points of (partial) agreement and of disagreement. Statements by Lee and Nelder are in *italics*; my responses and comments are in Roman.

### 2.1 Modeling Strategies

Lee and Nelder write, “*However, we believe that such a choice is inappropriate because the choice of an estimation method for a particular parameterization (marginal parameter) should not pre-empt the process of model selection.*” I agree. Estimation methods are a means to an end and usually not, themselves, the end (in methods research they can be the goal). Of course, the estimation method might influence model choice in that an inefficient method may miss important covariates and an inappropriate method may lead to bias. Sometimes the means/ends distinction gets blurred. For example, several years ago someone wrote to let

me know that he thought the EM estimate was the absolute best; far better than the MLE!

*Unified Probability Models are absolutely necessary:* I do take issue with this claim. One should not discount the effectiveness of analyses and algorithms that are not fully probability-based or comprehensive. These have and will continue to play an important role. While a unified approach with marginals, conditionals, etc., all generated by a joint distribution is without question the ideal, often it is not attainable. Data limitations, limitations in scientific understanding and computing constraints can thwart use of this holy grail. Even attainment can be illusory because the unified model may not be correct and may mislead. So, while I favor the unified approach, I’m very comfortable with an approach that validly and effectively addresses a specific goal.

“...so that care is necessary in making inferences about unobservables.” Absolutely! Extreme care and caution are most definitely needed. Inferences on latent effects are always model-based to some degree, and some assumptions cannot be verified empirically. For example, models using the standard Poisson distribution as baseline rather than the more general negative binomial will “identify” unaccounted (extra-Poisson) variation and allocate it to a latent effect. If a negative binomial model is used, much of this variation will be absorbed into the baseline model. Both approaches can produce similar predictions of observable quantities, but will produce very different inference for latent effects. All modeling approaches need to deal with such issues, and the h-likelihood is not a panacea. In contrast, use of latent variable models and hierarchical models to generalize the mean and association structure of models for observeds is quite safe. Therefore, I agree with Lee and Nelder that focus on the prediction space rather than the parameter space avoids mis- or over-interpretation of parameter estimates.

### 2.2 H-likelihood and Competitors

“...that when applied appropriately h-likelihood methods are both valid and efficient in such settings.” It is most surely the case that in some settings, with

---

Thomas A. Louis is Professor, Department of Biostatistics, Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland, USA (e-mail: tlouis@jhsph.edu).

an appropriate parameterization, the h-likelihood approach is valid and efficient. However, it is not globally valid and even when it is valid may perform no better than, and possibly worse than other approaches.

“However, GEEs cannot (generally) be integrated to obtain a likelihood function [McCullagh and Nelder (1989)] and therefore may not have a probabilistic or likelihood basis.” True, but GEEs can be very effective, especially for population-targeted inferences. I agree with Lee and Nelder that likelihood-based approaches or likelihood-like (marginal, partial, profile,...) approaches should be used when available and their use is essentially necessary when making inferences on latent effects.

“HGLMs allow a synthesis of GLMs, random-effect models, and structured-dispersion models.” They do synthesize, but aren’t alone in accomplishing this task.

Bayes is like the Adjusted profile h-likelihood (APHL). Well, that’s one way to put it. The other way is that the APHL is like Bayes. Regarding extended Likelihood versus (empirical) Bayesian approaches, one can think of the h-likelihood as prior augmented likelihood, an attractive approach to stabilizing and smoothing MLEs. However, taking full advantage of the structure requires moving away from mode/curvature inferences and, at least for some non-standard goals, employing the fully Bayesian formalism.

*Poor performance of plug-in empirical Bayes (EB).* Yes, naive EB produces a too-low variance estimate (more generally, an incorrect shape and association structure), unless the estimates of prior parameters are very precise. This observation motivated the Laird/Louis bootstrap and Carlin and Gelfand’s matching approach. These have been supplanted by Bayes empirical Bayes (BEB) with a hyper-prior from which prior parameters are sampled. BEB has proven very effective in producing procedures with excellent frequentist (as well as Bayesian) properties. See, for example, Table 3.4 in Carlin and Louis (2000) and Table 5.6 in Carlin and Louis (2009).

*Priors and hyper-priors.* Lee and Nelder state, “In Bayesian analysis, priors can give information on unidentifiable model assumptions, so that it is hard to know whether the information is entirely coming from the uncheckable priors.” Yes, and ditto for modeling assumptions whatever the approach. Care is needed.

In Section 4.3.1, Lee and Nelder criticize use of  $\sigma^{-2} \sim \text{gamma}(0.0001, 0.0001)$ . The problems with using this prior and a  $\text{gamma}(\alpha, \alpha)$  more generally are

well known. Though the mean is 1 and coefficient of variation is large, most of the prior mass is in the interval (0, 1]. It’s better to use a uniform prior on  $\log \sigma$  in a bounded interval with the bounds selected to respect measurement units. It is most definitely the case that more research is needed on selecting hyper-priors that produce good frequentist properties. This and other examples highlight the need for sophistication and care when exploring the latent world.

### 2.3 Goals that Challenge the H-likelihood

*Accounting for uncertainty.* Lee and Nelder make the important point about the need to account for uncertainty, but can’t avoid “dissing” (empirical) Bayes. They state, “The h-likelihood approach takes into account the uncertainty in the estimation of random effects, so that inferences about unobservables are possible without resorting to an EB framework.” The h-likelihood may take this uncertainty into account, but it does not ensure that all relevant uncertainties migrate into the inferences. For example, it does not allow for adjusting the shape of or association structure in the distribution of random effects, whereas the Bayesian formalism introduces both of these along expanding the spread by integrating over the posterior hyper-prior.

*Nonstandard goals.* Regarding goals, while the h-likelihood and other purely likelihood-based approaches can be effective in making inferences on measures of central tendency and linear functions of target parameters, they have a difficult time in structuring an approach for nonstandard goals whereas the Bayesian formalism is successful. For example, consider estimating the ranks of the  $\theta_k$  in a two-stage model,  $[\theta_1, \dots, \theta_K]$  i.i.d.  $G$ ;  $[Y_k | \theta_k]$  i.n.d.  $f_k(y_k | \theta_k)$ . As detailed in Lin et al. (2006), if the  $\theta$ s were observed,  $R_k(\theta) = \sum_{v=0}^K I_{\{\theta_k \geq \theta_v\}}$ ;  $P_k = R_k / (K + 2)$  with the smallest  $\theta$  having rank 1. Ranks/percentiles that minimize posterior expected squared-error loss for the ranks are their posterior mean or a discretized version,

$$\bar{R}_k(\mathbf{Y}) = E[R_k(\theta) | \mathbf{Y}] = \sum_v \text{pr}[\theta_k \geq \theta_v | \mathbf{Y}],$$

$$\hat{R}_k = \text{rank}(\bar{R}); \quad \hat{P}_k = \hat{R}_k / (K + 1).$$

The model can be generalized to BEB and is effective in both Bayesian and frequentist evaluations. Similarly challenging inferential goals are handled well (if handled with care!) by the Bayesian formalism, including proper accounting for uncertainty.

*Computational challenges.* Lee and Nelder write, “However, the computation of the ML estimation of the parameters can be a complex task because of intractable integration.” Yes, finding the MLE and developing appropriate inferences can be complex, and expansions around the mode may not be up to the task. Markov chain Monte Carlo methods have enabled likelihood-based and Bayesian-based analyses of complex data and models. Use them, but carefully!

*Resorting to (empirical) Bayes.* It is strange that Lee and Nelder characterize use of empirical Bayes a “resort.” In this day and age is the Bayesian formalism to be avoided? Have the last 20–25 years passed Lee and Nelder by? Most statisticians have gone beyond the Bayes/frequentist polemic of the 1980s and early 1990s. Yes, there are challenges, but use of the Bayesian formalism in both its objective and informative-prior forms, burgeons. Its use is by no means a panacea, but carefully employed, it is very effective in addressing both Bayesian and frequentist goals.

### 3. SUMMARY

Lee and Nelder provide considerable food for thought, considerable light and some heat, heat produced by their over-promotion of h-likelihood. I support Lee and Nelder’s goal of attempting a unified analysis based on full probability modeling, but note that the Bayesian formalism is best suited to this task.

Use of the full probability calculus, empowered by modern computing, brings in (most) relevant uncertainties, produces properly shaped and calibrated confidence regions and enables addressing nonstandard goals such as ranking. However, I caution that full probability modeling isn’t always available or valid and in many situations compromises are necessary.

Whatever the approach to analysis, care, evaluation, and sophistication are needed, especially when structuring inferences for latent effects. Polemic and over-promotion distract from the important issues and goals. These should be replaced by aggressive scientific evaluations and energetic discourse.

### ACKNOWLEDGMENT

Support provided by Grant R01 DK061662, from the U.S. NIH, National Institute of Diabetes, Digestive and Kidney Diseases.

### REFERENCES

- CARLIN, B. P. and LOUIS, T. A. (2000). *Bayes and Empirical Bayes Methods for Data Analysis*, 2nd ed. Chapman and Hall/CRC Press, Boca Raton, FL. [MR1427749](#)
- CARLIN, B. P. and LOUIS, T. A. (2009). *Bayesian Methods for Data Analysis*, 3rd ed. Chapman and Hall/CRC Press, Boca Raton, FL. [MR2442364](#)
- LIN, R., LOUIS, T. A., PADDOCK, S. M. and RIDGEWAY, G. (2006). Loss function based ranking in two-stage, hierarchical models. *Bayesian Anal.* **1** 915–946. [MR2282211](#)